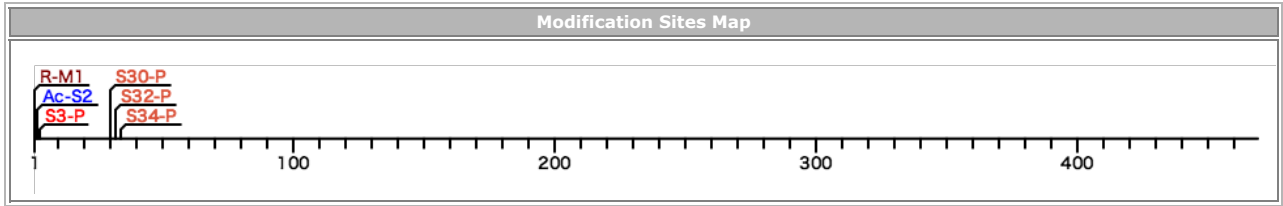


ID	Accession	GeneName	Chr.No.		Description
2A5E_HUMAN	Q16537	PPP2R5E	14q23.2	63838075..64010092	Serine/threonine-protein phosphatase 2A 56 kDa regulatory subunit epsilon isoform



Click a modification site to display information in detail.

Site no	Amino acid	Type	Division	Detail
3	S	P	Lab	140320_tita_C18_.mgf[F017426]
3	S	P	Lab	140320_OVISE_SCE_.mgf[F017431]

**Protein Sequence**

MSSAPTTPPS VDKVDGFSRK SVRKARQKR S QSSQFRSQG KPIELTPLPL LKDVPSSEQP ELFLKKLQQC CVIFDFMDTL SDLKMKEYKR STLNELVDYI TISRGC LTEQ TYPEVVRMVS CNIFRTLPPS DSNEFDPEED EPTLEASWPH LQLVYEFFIR FLESQEFQPS IAKKYIDQKF VLQELLEFD S EDPRERDYLK TVLHRIYGKF LGLR AFIRKQ INNIFLRFVY ETEHFNVAE LLEILGSIIN GFALPLKAEH KQFLVKVLI LHTVRSLSLF HAQLAYCIVQ FLEKDPSTE PVIRGLMKFW PKTCSQKEVM FLG ELEEILD VIEPSQFVKI QEPLFKQIAK CVSSPHFQVA ERALYWNNE YIMSLIEENS NVILPIMFSS LYRISKEHWN PAIVALVYV LKAFMEMNST MFDELTATYK S DRQREKKKE KEREELWKKL EDLELKRGLR RDGIIP

Backcolor of amino acid : Yellow -> site of modification, gray -> in front of processing